

## SEQUENCE LISTING

<110> Mount Sinai School of Medicine of NYU

<120> TRP8, A TRANSIENT RECEPTOR POTENTIAL CHANNEL EXPRESSED IN TASTE RECEPTOR CELL

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Pro Ala Leu Val Tyr Thr Asn Leu Ile Thr Phe Ser Glu Glu Ala Pro
                               665
Leu Arg Thr Gly Leu Glu Asp Leu Gln Asp Leu Asp Ser Leu Asp Thr
                           680
Glu Lys Ser Pro Leu Tyr Gly Leu Gln Ser Arg Val Glu Glu Leu Val
                        695
                                           700
Glu Ala Pro Arg Ala Gln Gly Asp Arg Gly Pro Arg Ala Val Phe Leu
                                       715
                   710
Leu Thr Arg Trp Arg Lys Phe Trp Gly Ala Pro Val Thr Val Phe Leu
                                    730
               725
Gly Asn Val Val Met Tyr Phe Ala Phe Leu Phe Leu Phe Thr Tyr Val
                                745
           740
Leu Leu Val Asp Phe Arg Pro Pro Pro Gln Gly Pro Ser Gly Pro Glu
                           760
Val Thr Leu Tyr Phe Trp Val Phe Thr Leu Val Leu Glu Glu Ile Arg
                       775
                                           780
Gln Gly Phe Phe Thr Asp Glu Asp Thr His Leu Val Lys Lys Phe Thr
                   790
                                       795
Leu Tyr Val Gly Asp Asn Trp Asn Lys Cys Asp Met Val Ala Ile Phe
                                   810
               805
Leu Phe Ile Val Gly Val Thr Cys Arg Met Leu Pro Ser Ala Phe Glu
                               825
           820
Ala Gly Arg Thr Val Leu Ala Met Asp Phe Met Val Phe Thr Leu Arg
                           840
                                               845
Leu Ile His Ile Phe Ala Ile His Lys Gln Leu Gly Pro Lys Ile Ile
                       855
                                           860
Val Val Glu Arg Met Met Lys Asp Val Phe Phe Phe Leu Phe Phe Leu
                   870
                                       875
Ser Val Trp Leu Val Ala Tyr Gly Val Thr Thr Gln Ala Leu Leu His
               885
                                   890
Pro His Asp Gly Arg Leu Glu Trp Ile Phe Arg Arg Val Leu Tyr Arg
                                905
Pro Tyr Leu Gln Ile Phe Gly Gln Ile Pro Leu Asp Glu Ile Asp Glu
                           920
                                                925
Ala Arg Val Asn Cys Ser Thr His Pro Leu Leu Glu Asp Ser Pro
                        935
Ser Cys Pro Ser Leu Tyr Ala Asn Trp Leu Val Ile Leu Leu Val
                    950
                                       955
Thr Phe Leu Leu Val Thr Asn Val Leu Leu Met Asn Leu Leu Ile Ala
                                    970
Met Phe Ser Tyr Thr Phe Gln Val Gln Gly Asn Ala Asp Met Phe
                                985
           980
Trp Lys Phe Gln Arg Tyr Asn Leu Ile Val Glu Tyr His Glu Arg Pro
                           1000
Ala Leu Ala Pro Pro Phe Ile Leu Leu Ser His Leu Ser Leu Thr Leu
                                           1020
                       1015
Arg Arg Val Phe Lys Lys Glu Ala Glu His Lys Arg Glu His Leu Glu
1025
                   1030
                                       1035
Arg Asp Leu Pro Asp Pro Leu Asp Gln Lys Val Val Thr Trp Glu Thr
                1045
                                    1050
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Val Gln Lys Glu Asn Phe Leu Ser Lys Met Glu Lys Arg Arg Arg Asp 1060 1065 Ser Glu Gly Glu Val Leu Arg Lys Thr Ala His Arg Val Asp Phe Ile 1080 1085 1075 Ala Lys Tyr Leu Gly Gly Leu Arg Glu Gln Glu Lys Arg Ile Lys Cys 1095 1100 Leu Glu Ser Gln Ile Asn Tyr Cys Ser Val Leu Val Ser Ser Val Ala 1115 1110 Asp Val Leu Ala Gln Gly Gly Gly Pro Arg Ser Ser Gln His Cys Gly 1130 1135 1125 Glu Gly Ser Gln Leu Val Ala Ala Asp His Arg Gly Gly Leu Asp Gly 1140 1145 1150 Trp Glu Gln Pro Gly Ala Gly Gln Pro Pro Ser Asp Thr 1155 1160 1165